

IN THE SPECIFICATION

Kindly enter an amendment to the specification.

Page 11, replace the paragraph starting at line 28 with the following:

The internationally recognized schemes for the classification and nomenclature of all enzymes from IUMB include proteases. The updated IUMB text for protease EC numbers are can be found on [[at]] the internet site: <http://www.chem.qmw.ac.uk/iubmb/enzyme/EC3/4/11/>. In this system enzymes are defined by the fact that they catalyse a single reaction. The system categorises the proteases into endo- and exoproteases. Endoproteases are those enzymes that hydrolyse internal peptide bonds, exoproteases hydrolyse peptide bonds adjacent to a terminal α -amino group ("aminopeptidases"), or a peptide bond between the terminal carboxyl group and the penultimate amino acid ("carboxypeptidases"). The endoproteases are divided into sub-subclasses on the basis of catalytic mechanism. There are sub-subclasses of serine endoproteases (EC 3.4.21), cysteine endoproteases (EC 3.4.22), aspartic endoproteases (EC 3.4.23), metalloendoproteases (EC 3.4.24) and threonine endoproteases (EC 3.4.25).